

Phe Leu Met Lys Thr Asp Gln Leu Pro Phe Glu Lys Ala Tyr Glu Lys
130 135 140

Leu Gln Ile Leu Lys Pro Glu Ala Lys Met Asn Glu Gly Phe Glu Trp
145 150 155 160

Gln Leu Lys Leu Tyr Gln Ala Met Gly Tyr Glu Val Asp Thr Ser Ser
165 170 175

Ala Ile Tyr Lys Gln Tyr Arg Leu Gln Lys Val Thr Glu Lys Tyr Pro
180 185 190

Glu Leu Gln Asn Leu Pro Gln Glu Leu Phe Ala Val Asp Pro Thr Thr
195 200 205

Val Ser Gln Gly Leu Lys Asp Glu Val Leu Tyr Lys Cys Arg Lys Cys
210 215 220

Arg Arg Ser Leu Phe Arg Ser Ser Ser Ile Leu Asp His Arg Glu Gly
225 230 235 240

Ser Gly Pro Ile Ala Phe Ala His Lys Arg Met Thr Pro Ser Ser Met
245 250 255

Leu Thr Thr Gly Arg Gln Ala Gln Cys Thr Ser Tyr Phe Ile Glu Pro
260 265 270

Val Gln Trp Met Glu Ser Ala Leu Leu Gly Val Met Asp Gly Gln Leu
275 280 285

Leu Cys Pro Lys Cys Ser Ala Lys Leu Gly Ser Phe Asn Trp Tyr Gly
290 295 300

Glu Gln Cys Ser Cys Gly Arg Trp Ile Thr Pro Ala Phe Gln Ile His
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Lys Asn Arg Val Asp Glu Met Lys Ile Leu Pro Val Leu Gly Ser Gln
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Thr Gly Lys Ile
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<211> 687

<212> DNA

<213> Homo sapiens

<400> 25

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 ggctgtctgc ctggaggaag gatgtccctg cactgataca gaaggctggt ctttaccctt 660
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<210> 26
 <211> 176
 <212> PRT
 <213> Homo sapiens

<400> 26
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 35 40 45
 Val Asn Val Ser Arg Gln Gln Pro Gly Pro Arg Ala Pro Gly Val Ala
 50 55 60
 Glu Leu Arg Val Pro Val Phe Asp Asp Pro Ala Glu Asp Leu Leu Ala
 65 70 75 80
 His Leu Glu Pro Thr Cys Ala Ala Met Glu Ala Ala Val Arg Ala Gly
 85 90 95
 Gly Ala Cys Leu Val Tyr Cys Lys Asn Gly Arg Ser Gln Leu Gly Ala
 100 105 110
 Val Cys Thr Ala Tyr Leu Met Arg His Arg Gly Leu Ser Leu Ala Lys
 115 120 125
 Ala Phe Gln Met Val Lys Ser Ala Arg Pro Val Ala Glu Pro Asn Pro
 130 135 140
 Gly Phe Trp Ser Gln Leu Gln Lys Tyr Glu Glu Ala Leu Gln Ala Gln
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 Ser Cys Leu Gln Gly Glu Pro Pro Ala Leu Gly Leu Gly Pro Glu Ala
 165 170 175

<210> 27
 <211> 901
 <212> DNA
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<400> 27
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 agaaacatgg aaagatgccga gaattcatgt tgtggaagaa gtagagccga gcagtggggg 240
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